

**Amendment to the specification:**

Amend paragraph 0132 as follows:

**[0132] SEQ ID NO: 67**

~~DLDQVRHILRRSWRSLSGNCKLLRRRMPSLRRSSRLEVTHMFQITLHFYKSTSRGGPV~~  
~~PSFCSE~~ and in split-Trp<sup>180</sup> split-Trp<sup>198</sup>, split-Trp<sup>203</sup> and split-Trp<sup>204b</sup> with a peptide of 10 residues possessing the sequence SEQ ID NO: 68 ~~(E/Q)RWIWRSGT~~. It is assumed that N<sub>trp</sub> and C<sub>trp</sub> of these clones associate spontaneously without the help of interacting proteins. In split-Trp<sup>44</sup> and split-Trp<sup>204b</sup> the mutation Gly8Cys was introduced during the fragmentation procedure. However, the influence of this mutation seems to be of minor importance as the deletion of the first ten amino acids still allowed split-Trp<sup>77</sup> to complement tryptophan auxotrophy (FIGS. 2 and 3).